

<110> Frudakis, Tony

<120> Efficient Methods and Apparatus for High-Throughput Processing of Gene Sequence Data

<130> 0201-0001

<140> US 09/964,059

<141> 2001-09-26

<150> US 60/274,686

<151> 2000-03-08

<160> 239

<210> 1

<211> 480

<212> DNA

<213> Homo Sapiens

<400> 1

gaattctttc	cagaaggctt	tccatttact	tttcctagat	tcatacagaag	aatcattatc	60
tacagcagct	gtaactgatt	gaaatgtatt	ttatgaacaa	taagacttga	aagttaaaat	120
tgctccttta	tccatgtact	gaagaataaa	tattgtgaaa	gcagtcataa	aaacagaagt	180
aatcttttgg	tacctctgca	ttagaactct	ttattaacca	ggtgtattgc	cattcaacag	240
taatattttg	aaaggaatct	ctatTTTTga	gcagggtttc	acttctgctt	tttattttaa	300
acagtagact	tgaaatattc	agtaaccatg	ctataaagag	ctatgctgta	agacagcttt	360
ttctatttat	agagcatggg	tttgaaatta	taacaaagca	tgggttttat	cctgaaatca	420
ttcataaata	acacgtacca	aaactttaat	acgggctagc	cagtgtgagc	cagtgtgagc	480

<210> 2

<211> 22

<212> DNA

<213> Homo Sapiens

<400> 2

ttccagaagg	ctttccattt	ac	22
------------	------------	----	----

<210> 3

<211> 21

<212> DNA

<213> Homo Sapiens

<400> 3

gttcaagtct	tggtacgtgc	t	21
------------	------------	---	----

<210> 4

<211> 780

<212> DNA

<213> Homo Sapiens

<400> 4

gaattctttc	cagaaggctt	tccatttact	tttcctagat	tcatacagaag	aatcattatc	60
tacagcagct	gtaactgatt	gaaatctatt	ttatgaacaa	taagacttga	aagttaaaat	120
tgctccttta	tccatgtact	gaagaataaa	tattgtgaaa	gcagtcataa	aaacagaagt	180
aatcttttgg	tacctctgca	ttagaactct	ttattaacca	ggtgtattgc	cattcaacag	240
taatattttg	aaaggaatct	ctatTTTTga	gcagggtttc	acttctgctt	tttattttaa	300
acagtagact	tgaaatattc	agtaaccatg	ctataaagag	ctatgctgta	agacagcttt	360
ttctatttat	agagcatggg	tttgaaatta	taacaaagca	tgggttttat	cctgaaatca	420
tgctccttta	tccatgtact	gaagaataaa	tattgtgaaa	gcagtcataa	aaacagaagt	480
aatcttttgg	tacctctgca	ttagaactct	ttattaacca	ggtgtattgc	cattcaacag	540
taatattttg	aaaggaatct	ctatTTTTga	gcagggtttc	acttctgctt	tttattttaa	600

Volume in drive D:\ is
Directory of D:\

seq_listing.txt

54 KB

4/4/02

1 file(s)

Total filesize 54 KB

0 kilobytes free